STIC Biotechnology Systems Branch

RAW SEQUENCE LISTING
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Application Serial Number: _ *10/522,592* _____

Source: _____ *PCT/10* __

Date Processed by STIC: _____ *2/22/05* _____

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Revised 01/24/05

PCT

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/522,592

DATE: 02/22/2005
TIME: 13:27:10

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Output Set: N:\CRF4\02222005\J522592.raw

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    4         BERTHOMME, Herv,
    6  <120> TITLE OF INVENTION: Novel method for analyzing nucleic acid and use thereof
    7         for evaluating the degree of mRNA editing of the
    8         serotonin 5-HT2C receptor
   10  <130> FILE REFERENCE: 347153
C--> 12 <140> CURRENT APPLICATION NUMBER: US/10/522,592
C--> 12 <141> CURRENT FILING DATE: 2005-01-25
   12  <150> PRIOR APPLICATION NUMBER: PCT/FR 2003/002 339
   13  <151> PRIOR FILING DATE: 2003-07-24
   15  <150> PRIOR APPLICATION NUMBER: FR 02/09 524
   16  <151> PRIOR FILING DATE: 2002-07-26
   18  <160> NUMBER OF SEQ ID NOS: 37
   20  <170> SOFTWARE: PatentIn Ver. 2.1
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pp 1,3

Does Not Comply
Corrected Diskette Needc.

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  847  <223> OTHER INFORMATION: Description of the artificial sequence: DNA derived
  848         from the mRNA coding for human 5-HT2C receptor              use
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RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/522,592

DATE: 02/22/2005
TIME: 13:27:10

Input Set : N:\DA\pto.da.txt
Output Set: N:\CRF4\02222005\J522592.raw

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SEQUENCES LISTING

<110> MADJAR, Jean-Jacques
      BERTHOMME, Herve *delete all foreign accent marks. The CRF*
      (or scientific symbols) software
      cannot
      process
      them.

<120> Novel method for analyzing nucleic acid and use thereof
      for evaluating the degree of mRNA editing of the
      serotonin 5-HT₂ receptor

do not use subscripts

The types of errors shown exist throughout
the sequence Listing. Please check subsequent
sequences for similar errors.

The types of errors shown exist throughout
the Sequence Listing. Please check subsequent
sequences for similar errors.

Input Set : N:\DA\pto.da.txt
Output Set: N:\CRF4\02222005\J522592.raw

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